

#5

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of: GUEGLER et al.



Serial No. 09/749,589

Filed: December 28, 2000

For: ISOLATED HUMAN TRANSPORTER
PROTEINS, NUCLEIC ACID MOLECULES
ENCODING HUMAN TRANSPORTER
PROTEINS, AND USES THEREOF

Art Unit:

Examiner:

Atty. Docket: CL000861

**SUBMISSION OF SEQUENCE LISTING
UNDER 37 C.F.R. § 1.821(a)**

Honorable Commissioner of
Patents and Trademarks
Washington, D.C. 20231

Sir:

In compliance with 37 C.F.R. § 1.821(a), Applicants submit the Sequence Listing,
including the paper copy of the Sequence Listing and the computer readable copy of the
Sequence Listing.

In the Specification:

Please enter the Sequence Listing between the specification and the claims of the
above-identified application.

09749589-051001

REMARKS

In accordance with 37 C.F.R. § 1.821(f), the paper copy of the Sequence Listing and the computer readable copy of the Sequence Listing submitted herewith in the above application are the same.

In accordance with 37 C.F.R. § 1.821(g), this submission includes no new matter.

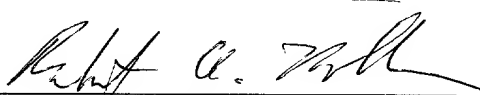
It is respectfully believed this application is now in condition for examination.

Early notice to this effect is earnestly solicited.

Respectfully submitted,

CELERA GENOMICS

By: _____



Robert A. Millman

Reg. No. 36,217

Date: May 10, 2001

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09749589-05400
T00T50-68564260



#5

SEQUENCE LISTING

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NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
AND USES THEREOF

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      20             25             30
Tyr Ala Leu Leu Leu Gln His Leu Pro Val Leu Val Trp Leu Pro Arg
      35             40             45
Tyr Pro Val Arg Asp Trp Leu Leu Gly Asp Leu Leu Ser Gly Leu Ser
      50             55             60
Val Ala Ile Met Gln Leu Pro Gln Gly Leu Ala Tyr Ala Leu Leu Ala
      65             70             75             80
Gly Leu Pro Pro Val Phe Gly Leu Tyr Ser Ser Phe Tyr Pro Val Phe
      85             90             95
Ile Tyr Phe Leu Phe Gly Thr Ser Arg His Ile Ser Val Gly Thr Phe
      100            105            110
Ala Val Met Ser Val Met Val Gly Ser Val Thr Glu Ser Leu Ala Pro
      115            120            125
Gln Ala Leu Asn Asp Ser Met Ile Asn Glu Thr Ala Arg Asp Ala Ala
      130            135            140
Arg Val Gln Val Ala Ser Thr Leu Ser Val Leu Val Gly Leu Phe Gln
      145            150            155            160
Val Gly Leu Gly Leu Ile His Phe Gly Phe Val Val Thr Tyr Leu Ser
      165            170            175
Glu Pro Leu Val Arg Gly Tyr Thr Thr Ala Ala Ala Val Gln Val Phe
      180            185            190
Val Ser Gln Leu Lys Tyr Val Phe Gly Leu His Leu Ser Ser His Ser
      195            200            205
Gly Pro Leu Ser Leu Ile Tyr Thr Val Leu Glu Val Cys Trp Lys Leu
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Pro Gln Ser Lys Val Gly Thr Val Val Thr Ala Ala Val Ala Gly Val
      225            230            235            240
Val Leu Val Val Val Lys Leu Leu Asn Asp Lys Leu Gln Gln Gln Leu
      245            250            255
Pro Met Pro Ile Pro Gly Glu Leu Leu Thr Leu Ile Gly Ala Thr Gly
      260            265            270
Ile Ser Tyr Gly Met Gly Leu Lys His Arg Phe Glu Val Asp Val Val

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Tyr Arg Val Asp Ser Asn Gln Glu Leu Val Ala Leu Gly Leu Ser Asn		
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